

FIG. 1A

FIG. 1B

FIG. 1C

FIG. 1



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Anopheles gambiae odorant receptor 1 genomic sequence (SEQ ID NO: 9)

Features:

- 1) Presumed Untranslated 5' and 3' regions are underlined.
 - 2) Potential TATA box transcription initiation signal is double underlined.
 - 3) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
 - 4) Introns are tentatively assigned and are shown in lower case.
- Exons are boxed.

AGCTTTGTTCAATTATGTTGAAATCTAGCCCATTTTGTAAGTGCTGAACGACGAAGAACAATACGAAAGTACCTCGT
CCGAACACTATCAACATTAAATTATACCAAGCTAGAAGAAGATAATTATAGTCAAGCCTCAACATCATAGGAAACTTT
AGCAAAACCATTTAATTTACATGATGATAAGTCCCACCTCTTACCCAGCACAGGTTTGAGAAGGACGAAAGTATCT
TTACGATAAATATTACTCTAAGGTAGTTTGAATAAATAAATAATTACGTGCAAGTGGTGCAATCGGACATCATTC
GAAAGAATCTACTAAGTCATACACACACCCAGACCGACGTAGTTTCATCTAGAAAAAACGGTCAGCTCCATC
GAACACGTCAGGACATAACTGCGACATGCGTATGGTCAGTTCCTACTAGTGCCAACACTGGTTCAGGGCACTACCTT
CCGAAGCAGTAGAACCTAATGTATTGGAAATTATTAGGACATACTGCAACATGCATATGGCTAGTTCGGCTGGTACC
AACGATGGCACCCAGGACACTATCTGCGGCCCTTGTAATAATCACTGTAAAAATCTATACAAAAACGGCTTTACCCATACT
TTATCACAAAAACGGCAGGTGAGGGCTGGATTGCTTCAAAGCATTAGAAATATATAATTTCAAAGTCCATAATCTCC
TTAAAGATAGACaCAGTAGAGAACACATTTAGTGCTCTTTTCGTCGAGTTAGTTGCCCTTCTCAAGTAAGCGTTT
AATGCTCAATTGTTGTAGATTTCGTTGGATGACTCTCGCTACGTGCTATAGTGGTCAATACTTCCAATTAGATTTCAT
AATTAGTTTCCAATTGTCCACGGAAACCCaCAAAAGAAAAAAACTTGATCTAGGGTGGAAATTTTCGAGAACa
ATTGGACACTTCAT**ATG**AAAAAGGACAGCTTTTTCAAAAATGTTAAATAAACACCGTTGGATCCTTT]gttggatttca
attctccaaattctgcagaataattctgcaaatTTTacaaaactgctcaaccacaataattccaattaatcatctg

FIG. 1A

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aacatttaaaactgataattaagatgagtaattgcttcgtcatcacctaagaaatcgattagtttggataaaaagaa
caaattgaaatacaataaagtcctgaattttattcgaataacggcttgaaactcatttatttcaaaaacctttgaga
aattcctcgttgaaaattgggtcctctatagtttctgtaaacgggccacttcaaaagcaagaactaacaataatcataat
tatgggtgcaagtaactatcagtaaccagtaatcgccattaaaaacttttctcaatttgcggtcgttacccggctaaa
tacagagcagagtaacgggaagtgatcaacgtcgctattagataacgaggaacccctccgaagggtgtgtgaagg
accttttcaaatgaaaccaagtactgtttccagttttaaatggatagttataaaatgagccgttcaacgatcggg
catcatttgagtttcatcttcgaggagaaaatagatcagtgccactgtttaaccgaaaagtaatgaagctgaacaaact
gaaccacgggtgggatgcgtacgatcgacgggatcgttctggttgcatgtgttgaataatttagG CCTAT
GGCCACCGGAAGATACGGATCAGGCAACGCGTACATCGCGTACGGTTGGGCTTTGCGGATCATGTTTCTA
CATCTGTACGCTCTAACGCAAGCCCTATATACTTCAAGgATGTGAAGGATATTAATgtgagtcctctagttagctattag
tgttccacctgtccataaatctgtcttttattgggtadGACATCGCAAAATGCATTTGTTCTGCTTATGACTCAAGTGA
CGTTGATCTACAAGCTGGAAAAGTTTAACATAACAATCGCACGGATTCAGGCTTGCTCGCGCAAGCTTAACCTGCACA
CTGTATCACCCGAAACAGCGCGAAGAAATTCAGgtaagcctgctgggaaatatgactaaaaagagtgctaaacaaacga
ctctcctccaaatgtadCCCCGTTTTACAATCGATGAGTGGAGTGTTTTGGCTGATGATCTTTCTCATGTTTGTGGC
TATCTTCACCATCATCATGTGGTTATGTCGCCAGCCTTCGACAAATGAACGTCGTCGCCcGTGCCGGCCTGGTTCC
CGGTGGACTATCACCATTCGGACATAGTGTACGGTGTACTGTTCTGTATCAAACCATTTGGAATCGTCATGAGCGCA
ACGTACAACCTTCTCGACCGATAACCATGTTTTCGGGCTTGATGCTACACATAAAATGGACAAATTTGTGCGGCTTGGTAG
TATGGTTAAAAAGgtgagttacggcgactacttgccctccagtaaggacaggagtttggttccggttatgatcatt
ttatcadCTTGGACATGACGTCCCTCCCGAACGCCCAATTGGTCGCAACGGATGCGGAATGGAAAGAGATGCGAAAGC
GCATCGACCATCACTCCAAGGTGTACGGTACGATGTACGCTAAAGTAACGGAGTGTGTGCTGTTTCACAAGGACATC
TTAAGgtacgaattgggccaattaatgtgtcatttaaaaagcttgacccaactttcacagcttcggcgatgaagt
gcaggacattttccaaagGATCTATCTTCGGCGCAAGTATGCGCGTCTGTAAATTATCATTTGTATGACACTGCTGCAAC
TACCGGGGgCGATGTTACGATgGCCGATCTGCTGGGCTGTGGGTCTATTTGCTAGTAAaGACATCGCAAGTGTTTA

FIG. 1B

TTTCTGTTACGTAGGAATGAAATCTCCTATACGgtaggttggacacgtagaggaattaaatgtttgggaagaata
tcaataccaaatagtagtggttgcgttacagACGGATAAAATTACAGAGTTTGTGGGTTTCCAACTACTTCAAG
TTCGATAAGCGTACCAGCCAAGCAATGATATTTTCTGCAAAAtgtgagatagcggtgatattgtgcagtcagtaca
ttaaatacgttctctatttcagGACTCTTAAAGATGTTCACATCAAGGTGGGAAGTGCTTTGAAGGTTACGCTAAAT
CTTCACACATTTTTCAGgtatgtaattatgctgtggtatttagcttgaaataagctacaaactttgaaagtaattt
caatctgttttgaagATTATGAAGCTATCGTACTCCTATCTGGCCGTACTTCAGAGCATGGAAATCAGAGTAATGGtG
tTAATATCCctTAATTGTTGAAATTATATTTTGTAGATTTATGTCATAAAAGTAaTaTTTAATTTTATACATCAAACGT
AAGCCCGctaGTTTTCAAATTAGCCCTTTTCCAAAATTATCAAATTGATTTCGAATTGCAGAGTTTCAGGAATT
TAATCTGATAGGATATCTTGTTTATCCAATAGAGGTGTGGAAGCGTTCCCAAGCCATTTCGTTTGATAGTTTATAGCA
CCGTCGAGCAGTTGATCGCTGTGATCGCTAGGCGCACCTGATTTTATCTTATCTCGCACCTGTTATGGCAAGGGCG
CTTTTCACACGTTTCACACAATAATAATGCACATGTATAATGCATTCTTACTTTAGCATTTTGTACATATAATACC
AAAATTATGCATTTTATTTCTCACGCAACGATTAGAGGATGACTTcACAAAGGTCCATCTAGTGGTAGGAGGTATAC
AATTATACCTCTCAAAAATCTCACAGCataATGAGAAACAAAAGGATACCAAGCATACCCTTTTTTACTTGACAAATT
TCATTTGATTTATGTAATAAAGCACTGCaCGTCGACTTCCTAAAA

FIG. 1C

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FIG. 2A

FIG. 2B

FIG. 2C

FIG. 2

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Anopheles gambiae odorant receptor 2 genomic sequence (SEQ ID NO: 10)

Features:

- 1) Presumed Untranslated 5' and 3' regions are underlined.
- 2) Potential TATA box transcription initiation signal is double underlined.
- 3) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
- 4) Introns are tentatively assigned and are shown in lower case.
- 5) Exons are boxed.

GGGATCCTCTAGAGTCGACCTGCAGGCATGCAAGCTTCCCTCACCGTGACGTGCTAGAAATGGTTCAACATACTCGT
CCGGCAGAGCGAAGACGACGGAATGTCCCAGGAAATGTAATGAGATATCACAGCAAGTGAAACCCAAACCG
AGCTGTGCGCTTTGTGTGCGCTTTAAAAATGGCCCTTCCTTCGCCGATCTGCTTGGTTTCACACGCTTTCCCCAGG
AAATCCACTGACCACTGGCCACACATCAACCCGAGCGGAGCCTCAGTGCCCAGCGAAGCATATAATTTGCTCA
AAAAGTCACGGTACTCAATTAAATTGATTATAATCAATTTCGTGGCTTCCAACACACCCCTTCTTCCACAATCCATCG
CCGAGTGAGCGAGTATAAAGGTGAAGAAACGTACCTTGCGCTTGCTCACTAACTGAACCCGGATTTCAAAAGGAACA
TAAACCGCAACCCACAGCCGAAA**ATG**CTGATCGAAGAGTGTCGATAAATGGTGTC AATGTGCGAGTGTGGCTGTTC
TGGTCGTATCTGCGGCGCGGGTGTGCCGCTTCTGGTCGGCTGCATCCCGGTCGCCGTGCTGAACGTTTCCA
GTTCCCTGAAGCTGTACTCGTCCCTGGGCGACATGAGCGAGCTCATCATCAACGGATACTTTACCGTGCTGTACTTTA
ACCTCGTCGtaacgtgggcgaggggaggggcaataaaccttcccacttggtgataatttcataacctttccatgtgtt
ttttattctctgtttgttgccatccagCTCCGAACCTCCTTCTCTCGTGATCAATCGACGGAAATTGAGACATTTT
TTGAAGGCGTTGCCGCGAGTACGCTCTCCTCGAGgtaagtcattggttttctagtttttggggagtggttaca
ccataaccaccccgacggtaacatttgatcgtcccgcgaaaaatgtttgtacagAAAAATGACGACATCCGACCCCGT
GCTGGAGCGGTACACACGCGGGACGCATGCTATCGATAATCGAATCTGTGGCTCGGCCCTTCATTAGTGCCTGCT
TTGTGACCTATCCTCTGTTTGTGCCGGCGGCTACCGTACGGCGTCACGATACCGGGCGTGGACGTGCTGGCC
ACCCGACCTACCAAGTCTGTGTTGTGCTGCAGGTTTACCTTACCTTCCCGCCTGCTGCATGTACATCCCGTTCA

FIG. 2A

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CAGCTTCTACGGACCTGCACGCTGTTTGCGCTCGTCCAGATAGCGGCCCTAAAGCAACGGCTCGGACGCTTGGGGC
GCCACAGCGGCACGATGGCTTCGACCGGACACAGCGCCGGCAGCTGTTCCGCCGAGCTGAAGAGTGCTCTAAAGTAT
CACAAACAAATCATCCAGtaagtagacgctagtagactcgaccggattgccctccctcggggagggggttgcct
atttcgggatgcggcagcagcatcacacaaacggaaccattaatctccggtttcatgccgcacgggcact
gggtcatgttcacatccttccttccaaacacacacgcgcgtgcacgtacagATATGTTCATGATCTC
AACTCACTCGTCACCCATCTGTCTGCTGGAGTTCCTGTCTCGGGATGATGCTGTGCGCACTGCTGTTCTTGCT
AAGCATTGtaagtaaaatcgaccgacgtgcggtcgctagtcggtctccggactctcatttcgggactcaatcgttcc
atctctcaataadAGCAATCAGCTGGCACAGATGATAATGATTGGATCGTACATCTTCATGATACTCTCGCAGATGTT
TGCCTTCTATTGGCATGCGAACGAGGTACTGGAGCAGGtaatggcgctgaagctgagtttggttgagcgggttcgcta
tagatcggctgtcttacattgtgtgttctgcattggggttcgttttccctctccatttcagAGCCTAGGC
ATTGGCGATGCCATTACAAATGGAGCGTGCCCGACTTTGAGGAACCGATAAGGAACGGTTGATTCTAATTATTGC
ACGTGCTCAGCGACCGATGGTGGTAAGtttggctgatcgatgctctgttcaatgaacatggcacagaggctgtgta
aatagctgttcattaataagtttttcagaatgtatcgttttagtgatttaaacgcattgttctatgcaatggta
gcaacaatagaccgctttattaatccaagcttccttaggattgatttttaagagaaagataaaccatttt
tagtaaccaatttagttacaggaaaccaaatacacagaatttattattattattattattattattatt
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gttgtgtcttattattgtgtgttattattattattattattattattattattattattattattatt
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FIG. 2B

ttccagtaatccataataaaaaataaagtaataatagtaataatccagtaactgtagtaatac
acaataatctctaagaattaaaaattgcatcttgttaatgaaatatgttgattgttcgaatagttcagaaaaacttaaa
aatgcctcagcattaaacagtttttgaggttggttcagggcatttagtttagataatttagtattttaagcatttggtt
ttcattactacaaaaagcaaatttatgagtgaattactttcagttcttctaaacgcctatgtgtatgcaattacat
aacaatagctctctttttatttgcatcttttcccttagtaaatctcaaatctcttccctcttgca**ATTAAA**
GTCGGCAACGTGTACCCGATGACGTTGGAAATGTTTCAAAAATTGCTCAACGTCTCTACTCTTATTTACACTGCT
GCGCCGAGTGTAACAACTAA**ACTTAACCGGTAAACAACAATAATCCCCCTCATCACTATGCAAAAGACAGCAAGCAGCC**
GATCATCAAAACACCATTAGCAGCCACAAGTTACCAGCCGCTTATCCACGGGATTTGGTGGAAAGTTATTGCAC TG
AAGCTCTTTCACCCCAAATTTTCATGGAGGTTCCCTCTCAACCAACCCATTGAAGCGAATAAAAGTATCAGCAACCAG
GCGACGGTGAAAAAACGCTGCATTATTGTGCTTGCTTCAGCATTCAGCGAATGACTCTTAAACTTTTCCATTCAAA
AGTCGCGATGCTCACGATACGGAGCGGTGTGTTCGATCCGCCGAGTGCACTCGCAAGCCGGTGATGTTGCCGGT
GGAAATGCACAGATCGACACAGCGATAGATAATCGTTTGTTCGCGTAAATGGAGGGAATAAGTAAAGTGCCAGCT
ACTTCATTTCATGTTAATTGAAACTCAAGCCAAACATGCAGAACCCGGTTGGTGTGTCTCCGCTCCGGGA
AAGGTCCTCTGCTCCGGGCATGGATTCTTTCCCCCTCCGGTGGTGGGGTATTGTTTAGGTTTATTATTACAAA
TTCATATCCTTCCGCTTCCGCATCAGCCGACCCGGTGCGCCAGACAGATGTGCGGGCAACAATACTATGC
ACGAACATGGCCAACAACACAGCTTCTATCTGTGTCGCACTGTCTCGCTTTCCCGCTGCGTTGCTTGTA
GTACTATCATTTGTTTAGTCCACGGGTTTACTTCTAATTCCATTGCACCACGCAAAAGGCTCATCCTTTGCTCGTT
CCGGTTGCAACTTCGACAAGCGCATGGTTGGGATACGAACAACAATACTACTCCACCCTACTACTACTG
CCACCACCACTAACACACTACACTTGGTTGGAGCTTGCAAGACCCACAAGCAACACGATACAAGCTAGTAGCT
GCTGTGTGCGCTCGAGTCAGCCGACGGTACAAGGTTTAACCGGTACAAGCAACTCCCGGACCGATCCCAAACTCTG
ACAAGGCACGGGCCGATCCGGCAGTACGGTCGGAAACATGGAAATGTTTAAATAAAACTGTAAATTGTCAATCGC
TGCTACAAGTTGTGACACAGGGAGAGAGACAGAGCGCCGATGGTGATGGTGTAAGATAGATACAGGAA
AAGAGCGAGAAACATTGTACGATTGGTGTGTTAGCAAAATTGATTTCCACTGATTTTGAGTGCAAAATTAAATGC
ATCGAAAATTTGCCATTACGGGTAAGTTGCTCGTGACGGATCCCCGGCTGCAGGAATTCGATATCAAGCTTAT
CGATACCGTCGACCTCGAGGGGGCCCGGTACCCAGCTTTTGTTCCTTTTAGTGGA

FIG. 2C

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FIG. 3A

FIG. 3B

FIG. 3

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Anopheles gambiae odorant receptor 3 genomic sequence (SEQ ID NO: 11)

- Features:
- 1) Presumed Untranslated 5' and 3' regions are underlined.
 - 2) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
 - 3) Introns are tentatively assigned and are shown in lower case.
 - 4) Exons are boxed.

AAGCAGAACACATCAAGAAGCAATTAGGTGTGTCGTACGTTAGCAAGTAGTTCGCGAGGAGGAATAAAATAG**ATGCC**
TTCTGAGCGGCTTCGTCTCATTTACTTCCTTCGGAACCTCCTCAAGACAAACGCACGATGGTACTGCCAAAATTAAAGG
ATGAAACAGCAGTGATGCCGTTTCTGCTGCAAAATTCAAAACCATTGCCGGACTGTGGGTGACCGTTCCCGAGCGGTAC
CGTTTTATCTCATCTTTTCCCTACTTCTGCGCGATGGTGTCTACCCAAAGTGCTGTTCGGTTATCCAGATCTCGA
GGTTGCGGTACGCGCACGGCCGAGCTGATGTTCGAATCGAACGCATTTCTTCGGCATGCTAATGTTTCCTTTCAAC
GCGACAACTACGAGCGGATTTGGTGCATACGCTGCAGGATCTGGCAGCTCTAGgtgagtatgcagccaatcgattgttc
caaaccttcgcaacatccttcgtaacactgctacactttcagTCCTCCAAGACCTACCCACAGAGCTGGGAGAGTAC
CTGATCTCAGTGAAACCGACGGGTCGATCGGTTCTCCAATAATTACTGCTGCTGTCACTTTTCCATGGCAACGTTCTT
TTGGTTCATGCCCGTCTGGACGACCTATTCCGCCTACTTTGCTGTGCGCAACAGCACGCCGGAACCGGTCGAGCACGTGT
TGCACCTCGAGGAAGAGCTGTACTTCCTGAACATTCGGACTTCGATGGCGCACTATACGTTTATGTGGCCATTATG
TGGCCACGATCTATACGCTCGGGTTTACCGGTGGCACAAAGCTGCTGACCATTTTCAGCAATGTTAAGTACTGTTC
GGCCATGCTGAAGCTCGTTGCACTCCGAATCCACTGTCTAGCGAGAGTAGCGCAAGACCGAGCGGAAAGGAGCTGA
ACGAGATTATTCCATGCATCAGCGGGTACTCAAgtaagtaaatcaaatgaaagtttgcagggaataaacttgag
tgtgtctgacccgtgcacatcctagCTGCGTGTTCCTGCTGGAGACGACATTCCGCTGGGTATTTTTCGTGCAGTTC
ATTCAGTGTACAAATGATCTGGTGCAGTCTCATCCTCTACATAGCGGTGACGgtaatagcattttcgtcatttcgtta
GccttattcaatccatttttgtgaacgtgaatttccccagGGTTCAGCTCGACGGTAGCGAATGTATGTGTCCAG
ATCATTTTGGTGACGGTGGAAACTTACGGCTACGGCTACTTCGGAACAGATCTAACCCAGGAGGTGCTTTGGgtacc

FIG. 3A

ctttggatgaagcttcaaaaagtaattccaaattctgttttcgatttttcccccttttccactaaAGCTATGGCGTTG
CCCTCGCCATTTACGATAGCGAGTGGTACAAAGTTTCCATTTCGATGCGCGCAAACTTCGACTGCTACTGCAACGA
TCCCAAAACCGCTCGGCGTAACGGCGGAAAGTTTCGCTTCGTCAATGTGCCCCAGTTTGGCAAGgtaacattaat
tacagtttgaaaattctgaagaatgcatcttacttgcccttactgtgtgtccagATGCTCAAGATGTCCCTATTTCATT
TTACGTAGTACTGAAGGAGCAGTTT**TAG**GAGCTGCTGTTTCCACCCCTGGAAATGGCCCTTTTCGCACTGTCTTCTGT
TTGTTGGACGCACGCAGCACCGAGAGCGCCCTGCACGCACTGACGTATTTTGGCTACTTTGACGTTTGACCTTTG
ACAGCTGAAGGACAGGGTACAAATTTTGCTGCTGTATTACGCGCAGCGCATTTGGATACGAAACATTTGGCCACAAG
TTCTACGATTTTAGCGTTTATTACTGTTTCGTAGCAGCTTTTCCaCAATAAACACACACAATAACGTACCGACAG
TATTCTTTTCATTGTAGGATAGAGAGCCCGCCAGCAGCCAAACGCGCCGCAAAACGAAAGCGGCACCAACCG
GGGGAACACACGGAGCAAAACGAGAACAGACAGTAAACAAACCGGCCGGAACAAACACCGGTGCCCGGAA
ACGA

FIG. 3B

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FIG. 4A

FIG. 4B

FIG. 4

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Anopheles gambiae odorant receptor 4 genomic sequence (SEQ ID NO: 12)

Features:

- 1) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
- 2) Introns are tentatively assigned and are shown in lower case.

GGGGAAC TCCCCAC CCGACGAC GGAAGCTA ACGATGT GCAATTGA ATAGTCATTAGT
AGCGTTT TGTCTCGC AAACGAACTA ACCCTTTG ACTTTTAA GTTCACTA CCGTGAGGACAAAA
TCAATAA ATTAAATC GAGACCGT TGATGAGCA AAGAAAAA ATATTTTA CTGATTTTCATT
CGTTCCAT CGACTACA TAAATCAT AATTATAT GCCACATTTA TATAAG TTTTGTAT CATTTTTA
AACAAACA AAAATGCA TCCTTTCG AATATTAG TCAGGTTG TATCAACA **ATG**AAGTTT GAAGCTGT
TTCAAAAAT ATTCCTC CCCGACAC GGTCTTAT CCTTCGTG CTAAGGCT TTTTGCA TATCGTGGC
ATGAATGG GGCAGGAT TTCGGTC GCGAATTC GAGTTG GTGGCAT TTTTCTGT TCTATT TTAATCTT
TCTTGTAAT ACCGCCAC TAAACGG GGTACAC CGATGGT CACCAGC GTGTACG CACCAGTGTG
GAATTCCT GTTTAAT TGGCAAT ATTTACG GCGGCA GTATGT TCTTTG CCTACG ATGTGGCCACTTT
CCAAGCGT TCATCCAG GAAC TGAAGAG CCTTTTC GGTTTTGG taataatta aattcggttattgcat
catcattgt tctcttgca gTATGCTC ACATTCGT ACAGACTA AAGTATA AGCTGA CCGGTTCA ACCGTC
GAGCGGAT ATATCGC CCAAGT GCAACG ACCTGCA TGGGTG CTGTAA CGCTTTTCTACTGGAT
TGCACCGAT ACCTTCCA TCTGTGC GCACTA CTACAGGTC GACCAAT TCCACCGA ACCCGTGCGG
TTTGTGCA ACATTTAG AGGTGA AGTTCTA TTGGCTC GAGAA TCGCACCT CAGTCG AGGACTACAT
AACCTTCG TGTGATCAT GCTACCCG TCGTGTTAT GTGTGGT TACGTATG CAAATT TGAAGGTGA
TGACCATCT GCTGCAG CATTTGGAC ACTGTACACT GTACACCA GGATGACT ATAGAGATGGTAGA
GCAGTTGG AAAGCATG GCATCAG CGGAACG AACTGCC AGCCATAC GCAACGTG GGCAGAT
GCACAGTGG TTTACTG AAAATGC ATTAGGCT TTTGAACA CAGTCAATCC GATCGATG CTGATGCTGC

FIG. 4A

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ACTGGTTGACCTGCGTGTAAACTGGAGCATTTCTCTCATCTATCTAACGAACGTGgtagtttgtctt
gttggaatccaaaagaagatggctataattgaacttctattacagGGCATCTCGCTACAAATCGGTTACCGTGGT
GGTAAATGTTTCTTGCCACTGCGGAAACTTTCCTGTATTGTTACTTGGGACGCGCTTGCGA
CACAAACAGCAGCTGCTGGAGCACGCACTCTATGCTACACGGTGGTACAACTACCCAATAGCCTT
TCGCAGCAGCATTAGGATGATGTTGAGACAGTCGCAAGGCATGCACACATAACGGTGGGAAG
TTTTTTCGCGTTAATTGGAAGAAATTAGCAGGATTGTCAACTTATCCTACTCTGCTTACGTCGT
ACTTAAGGATGTAATAAAGATGGATGTACAG**TGA**ATGTMTMTMTTGGCTTGCAACGAATGA
AGTTTCCGAATCTATAATTAGATCTAGAAATTTAATCTAGATGTCATAATATGATCTTGGCCATGA
CCGGTTCCTGGTTTGGAAACCAATTCTCAAAACAATTTTGAACCTTAGGGCGAGGCATGAAATGTC
CCAAGAACCTATCCAAGTCTGGAACTACATATTACCGAATCTATCCCATTATTGCCCTCGGAACCT
GGTTGGTGCTAAATATTGTCCAAATGTGGTCCCTGACCTATCCAGACAAAGATCTTCAATTA
TTCCCTACCACTGGAACCTGATTAATTGATGTAGGAAGTCATGGAGGTGTTCAGGGAGAAATTTAAA
CACTAATGTTCCAACTCATTAATTCAAGGCCAATTCTATTTTTATATGCCCTACGGATTGATAC
GTATGTATTACTCCATTTCCTGGACTTTGTCTTATTCTGTGCTGATTTGGACGTGAAATGTTGA
GAAAAGATTCTTATTATGAGTGATACAGAGCCTTTAAATACTCCTACGTTGTGCTATTTTAA
GTATGGCCAGGCTAATCACAAATCGCTACTAATGAACAGAAATCTCTTAATTAAACCCCTTTCGAT
TGATAGTGTCAATGTGCGAGATAATTGAACCTGCAAAACgATACCTACCTTAAACGGAGCAG
AACACATCAAGAAGCAATTAGGTGTGTCGTACGTTAGCAAGTAGTTCCGAGAGGAATAAAAT
AG

FIG. 4B

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ANOPHELES GAMBIAE

Preferred DNA Codons

| Amino Acids | | Preferred Codons |
|---------------|-------|-------------------------|
| Alanine | Ala A | GCC GCG GCT GCA |
| Cysteine | Cys C | TGC TGT |
| Aspartic acid | Asp D | GAC GAT |
| Glutamic acid | Glu E | GAG GAA |
| Phenylalanine | Phe F | TTC TTT |
| Glycine | Gly G | GGC GGT GGA GGG |
| Histidine | His H | CAC CAT |
| Isoleucine | Ile I | ATC ATT ATA |
| Lysine | Lys K | AAG AAA |
| Leucine | Leu L | CTG CTC TTG CTT CTA TTA |
| Methionine | Met M | ATG |
| Asparagine | Asn N | AAC AAT |
| Proline | Pro P | CCG CCC CCA CCT |
| Glutamine | Gln Q | CAG CAA |
| Arginine | Arg R | CGC CGG CGT CGA AGA AGG |
| Serine | Ser S | TCG AGC TCC AGT TCT TCA |
| Threonine | Thr T | ACG ACC ACT ACA |
| Valine | Val V | GTG GTC GTT GTA |
| Tryptophan | Trp W | TGG |
| Tyrosine | Tyr Y | TAC TAT |

[http://www.kazusa.or.jp/codon/cgi-bin/showcodon\(con'd on next line\)
.cgi?species=Anopheles+gambiae+\[gbinv\]](http://www.kazusa.or.jp/codon/cgi-bin/showcodon(con'd on next line).cgi?species=Anopheles+gambiae+[gbinv])

FIG. 5

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| Name | SEQ ID NO |
|----------------------------------|---------------|
| Arrestin 1 (cDNA) | SEQ ID NO: 1 |
| Arrestin 1 (polypeptide) | SEQ ID NO: 2 |
| Odorant Receptor 1 (cDNA) | SEQ ID NO: 3 |
| Odorant Receptor 1 (polypeptide) | SEQ ID NO: 4 |
| Odorant Receptor 2 (cDNA) | SEQ ID NO: 5 |
| Odorant Receptor 2 (polypeptide) | SEQ ID NO: 6 |
| Odorant Receptor 3 (cDNA) | SEQ ID NO: 7 |
| Odorant Receptor 3 (polypeptide) | SEQ ID NO: 8 |
| Odorant Receptor 4 (cDNA) | SEQ ID NO: 13 |
| Odorant Receptor 4 (polypeptide) | SEQ ID NO: 14 |
| Odorant Receptor 5 (cDNA) | SEQ ID NO: 15 |
| Odorant Receptor 5 (polypeptide) | SEQ ID NO: 16 |
| Odorant Receptor 6 (cDNA) | SEQ ID NO: 17 |
| Odorant Receptor 6 (polypeptide) | SEQ ID NO: 18 |
| Odorant Receptor 7 (cDNA) | SEQ ID NO: 19 |
| Odorant Receptor 7 (polypeptide) | SEQ ID NO: 20 |

FIG. 6

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FIG. 7A

FIG. 7B

FIG. 7

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Anopheles gambiae odorant receptor 5 genomic sequence (SEQ ID NO: 21)

Predicted Exons: *ITALICIZED*, UNDERLINED AND **BOXED**
Introns: lowercase.

tctagacttgaaccatgacgggcattttattgagtcgttcgagttgacgactgtaccacgggaccaccggtttatcactactatt
aattaattataatagctttttagcagcctaccgggttttctcttgatatcttaagtcccatattgattatcaagatagaa
caacaactgtaccttaataatcattacgtacccttaatacaacctgtgcatcaaggagtttcggaaagcaaaaatccgattgct
gatgtgtcttgattccatccgattcgttactggttctgcaaaatcgccaataacggcaatgtccttategatgcttgaatcaacat
cacattgttgcatttcgtttttgcgtgcaaatatgttatttgcaagaaggcaagtatgtcttaagagtaatacaaatcgcgtg
tccatttttgcaccagtggtgccagaacccgtgccttttagctcttcgaatacatccgaccagtcagcaagtgcatd**ATGG**
TGCTACCGAAGCTGTCCGAACCGTACGCCGTGATGCCGCTTCTACTACGCCCTGCAGCG
TTTCGTTGGGCTGTGGGTGAACGACGCTATCGCTACAAGTTCCGGTTGGCATTTTA
AGCTTCTGTCTAGTAGTTATTCGGAAGGTTGCCCTTCGGCTATCCAGATTTAGAGAC
AATGGTTCGGGAACAGCTGAGCTGATTTTCGAATGGAACGTA**CTGTTTGGGATGTTG**
CTGTTTTCTCTCAAGCTAGACGACTATGATGATCTGGTGTACCGGTACAAGGACATATC
AAAGATTG^{gtgcgtgataatgattgataaaaggaaacctttgagcaactctatcccttcaag}**CTTCCCGTAAGGAC**
GTTCCCTCGCAGATGGGCGACTATCTGGTACGCATCAATCATCGTATCGATCGGTTTTC
CAAGATCTACTGCTGCAGCCATCTGTGTTGGCCATCTTCTACTGGGTGGCTCCTTCGT
CCAGCACCTACCTAGCCTACCTGGGGCACGAAACAGATCCGTCCTCCGGTCGAACATGT
GCTACACCTGGAGGAGGAGCTGTACTGGTTTCACACCCGCTCTCGCTGGTAGATTAC
TCCATATTCAACCGGCATCATGTGCTGCCCTACAATCTTTATGCTAGCGTACTTCGGTGGACT
AAAGCTGCTAACCATCTTCAGCAACGTGAAGTACTGTTCGGCAATGCTCAGGCTTGTG

FIG. 7A

GCGATGAGAAATCCAGTTCATGGACCCGGCTGGACGCGGAAAGGAACTGA
TCCGAAATCATCCGTCAATGCAATCAGAAAGCGCTAAAGtaagggtctgcccgggtatggtggataatacat
ctagtgccttcagATGTGTGGAGCTGTTGGAAATCATCTTTCGGTGGGTTTTCTGGGACAG
TTCATACAGTGCGTAATGATCTGGTGACGCTTGGTTCTGTACGTCCGCCGTTACGgtaacta
aaagcactgtagtctgtctgccacaccattcactgctgtctgtttgtcactcttcccagGGTCTCAGCACAAAG
CGGCCAAACCGTGGGTACTGTTTATACTGCTAACAGTGGAAACCTACGGATTCTGCTA
CTTGGCAGTGATCTTACCCTCGGAGGCAAGTTGTTATTCCGCTGAgtttcagttacttttccgttcccc
tctaaccgtaccactgtaccattgtttggagacagagcttgagcgtagCACGTGCTGCGTACGGTAGCCCTCTGG
TATCGCCGTTCCGGTTCGATTCAACGGGAAGCTTCGGAATGGTACTGCAGCGTGCCCCAGA
AACCGGTCGGCATCTCGGCTGGGAAGTTTGGCTTCGTCGACATTGAGCAGTTTGGGCAA
TgtatggggagaccttccactgtggcaagaagattttcttttattaatgcattcttttaattacagATGGCAAAACATCA
TACTCGTTCTACATCCTGTAAGGATCAATTTTAaggggaactccccaccaccagacgacggaa
agctaacgatgtgcaattgaatagtcattagtcgtttttgctcgcaaacgaaactaacctttgactttttaagttaactacggtgag
gacaaaatacaataaatacagaccggttgatgagcaaaagaaaaataattttactgattttcatttcgttccatcgacta
cataataataatgcccacattttattataagttttg

FIG. 7B

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Anopheles gambiae odorant receptor 6 partial genomic sequence (SEQ ID NO: 22)

These are the predicted last three exons of another candidate *Anopheles gambiae* odorant receptor.

Predicted Exons: *ITALICIZED*, UNDERLINED AND **BOXED**.
Introns: lowercase.

aacaccattatcgcaaaattagattaccggttgaaagcggtcccttcctggctgtttctcactctctctctctctctt
ttgatgccgtatgcgccgtgetatagctagTTATGCTTACCGGATGTTGCCGATCGCGCACCGTGCTTT
TCCGCATACGCCAGTGCACTTGATGGCGGTGGTGATGACGCTCTGCTGCCGACCGTT
TTCTGCTCGTGAGTCAGACCCTTTTCATTTCCTGCAATATCCTGTTTCTTCCCGACCCC
ACAGACGGTTAGACGGATATATGCTGGTAAAGTTTGTCTCTTCATGCTGTGCTTCTG
ATCGAGCTGCTGATGCTGTGCTGCTACGGTGAGGATATTGTGGAATCGgtaaggcaccaggc
ggtgatgagcgcgagtgagtaattgaagcttttgcctttaaacacatcagagCCTTGGGGTGATTGATGCCGCT
TACGGTTGCCGAATGGTACCGGGAAGGTCGGTGCGCTCCATCGATCCGTGCTGCAAA
TTATACACCGCAGCCAGTCCTCATACTGACCGCATGGAAATTGGCCCATCCAA
ATGAGTACTTTCAGTCAGgtgagttgccaatgattgccgtttggtaatttcagtaagagtgctctttcccttag
ATCCTGCAAGCTTCTGCTCTACTTTACCCCTCCTGAAGACCGTCTACGGGAATAAgtaa
gcgcgagagagagagagcagtatcgttcacccctttggatgaatcaatagatttctaatacgaaccattgaaaaatgaatca
acatttcgctagtgcacaatatgtaccattctatacagcttcaccacgaccaaagcgtttgtgcatcaggaccaaacacggttcga
caagccgcgtcacctgctggc

FIG. 8

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FIG. 9A

FIG. 9B

FIG. 9C

FIG. 9D

FIG. 9E

FIG. 9F

FIG. 9G

FIG. 9

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- $+$

 $+$ $+$

cctactagCCTGGGTTACGATAACATTTTCGGCGAGAGCGTCAAGACTGTGCTCGATAAG
GCAACCAACGAGACGTACACGGTGGATATACCCGGCTGCCCATCAAGTCTGTATC
CGTGGAATGCAATGAGCGGACCGGCTACATTTCTCTTCTATCTACCAGGTACGTTG
GCGGAATgtcctgcgcgtcacagttggcagtcagtgaggcaacacggcgaaatgggactaaaacgggtcttcacaga
gccaacacattcctacagcaattgcataaccttcggcggtcggactgggcaatgcagtacaacatcctcgcctaaagttagtcaat
tcgagcgacaaatgttgcgtgttagggctttttgtgataatagtcgtttttgtcctcctcgttatcaaaactctatcaacggaggaaa
tcattttcgtacaatgcctacagctcaagttcaaggtaatcgagcgggtgggatcaactttttattcattttgtctaacgccccca
tcaacaaattctatgttctcaatggcaaaagattactgccccaccaatcgccccaaacggcaaaagaaagcgacgattatga
agatgtccaaacattgccccgcgcgttatctgatgttgcggatggcttttactgtctgtactttcaggcacaaaaggaa
atgaaaccagcgagctcgtttgcggcttcggaggttcttcaggcactgagctgagtaactaaatcgacgatttttacgattc
tggatccagtttatgatgtggcctgcattacagtggaattataacctgatgttcatttcattgttaagtttgtgctggtaacg
ccgtaacgattaattctttcaaaagagattctttcaaaagagattcaaaatgtgtatacaaaatgctaaacgaatggaccgtacttg
agggttcggaaagtaacgttttaaaatattcatcaaatcctctgcaaaacttgtcttaattggtgcacaataagtttaaaact
gtggcggcagatgtgtcgtcgtccttccttcctccagcaagctcgtgcgaaataatttattccatcattttaatacagccgtttgtg
cattttaattagcaaaagcaataataaaagcagctaaccatccccattaaacaaagtcttcgggccaattgttatggcgtgga
aagtaatgggtttaccagtggaagtgtcctttcccatcgtgggtacttcgcgatatcttgtcttatacaagtgcatacagaaaaaa
ggacaaatcctccttgctatggtctaaggccagcttcggtaccgcttcggtatgcataaagtttgatgggtgttttaacatt
actccgcttaaccacctaattggacttttcatgttgactaaagttaaccagccaccagcggtacgaccgagccacggttgatt

FIG. 9B

FIG. 9C

FIG. 9D

 $+$

tttttgacgggtgtatagtaaattgcatactttaaggcgtgattttcaaatgtagcgttccgtatgcagaaacgccatggattatgc
aatttaacaatgctgtcttcttaacattcaaaataacggcttattaaggaactttttgtgcaatttggttttaacagcaaatagtttagc
tcagaacgatcacatttagtatcgcttcaacaagaactcttttaaacacacaatttgtaatgccattccctcgagaaagtttcttgctc
agtcctccttgcatcacagcaacaacaaacctgctcatgtttcctgctcgttttgaaacgttatttccgattcctgtgct
tgccccgttttcttacaatcaaccacaatggttcagatttcgctcttatttattgacccactgctttcgtgtaagccccgtggaaacaa
tgcgccaagctcagcatccagccatgcatgtataaatgagccacgacagatttttagacatcgctttcgtctgcacgggaggtggtt
ttattcttggttccgattccacgtccattcgtcctgggtccgcccgggaaacgtaagccgctgcggggaattacgcaatcgaa
aacgagccagaaaatgagcacgccaatgcaaaagaaaatcccccttttgagtggctcctgcacaaaacatgctccccggtgaagctgcgtatgt
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gtcttttagccgtgattgaattttctgaatatcacaggcgggcggtttgacctgcaaggttgcttccacacgagcattgcttt
ccgtaccgcggtgggcgagttttcaacgcacaaccttctacaagcaacgcacacaacgcctggagcgatatttaacagaaacaagaa
catcccgaacttcagcacatgccgtgatttgccctgttggaaaaagcttttgtagcgtgtgagttgaacgagctctattttccagcgat
gggtggcatttgtgtggcatgctatcgctcagcttttcttgaaatctttacctcctccattcgctccattagtacacgcgtatggaaaaatgg
gtgcaacggatcagaacggattttccgcgacagacttaataaagggaagcaacgcgttttttgcatgtgtagtgtttatgagcttt
atgccgttactttgcaattaaaaatagcaaaaaataaacagtttttttgtaagcggattacaaaagaatgtatcagaatattacgtg
aaacattcatttcagtgttaacgctcaaatagaaatagttttgtaacacggattgcataaccttgccggtatcggttacattttcgccta
acagtatgcaatctgttagctttgttgaatgactgcgttggttagacaatattttacacccggtaatattatctcacaatgtgc
aaaaaaatgtcaatctgtatcgattattcacaaaaatcagatccccggaaccagtgtagcccaatgtgctcttattgaattaccacga
acaaatcaacctgatccccgggtccgttggcaaacagcttgccccgaagccgctcagtttctgctgactaccgtgctgccattttgct
gccctcatcgaaacagataaacagaaggcaactcttgtgagcatcgcaatgccccgtctgaagttccgtcgaaaaatgggcctaattc
aatttgacgcatttaccgcgaacaattgcgcgaaggtgtcaagtggttccacgaactgcgacaacaagcacacacacaacac

FIG. 9E

aatgttatcgttttcggcatgtttctcgggtacaaagcgtgtggcgctatgtggcatgccgattcccagacagagtgatcgatagtaaa
tgtagccatatccggtagcattcaatttcttctatctcgcacaaagcccattctgggagcggtgtgaagctttcaaaagccat
tgtgaacaaatgtcctggttcggagggatgtcgggaaagcaaacacggtgccgccatcgctgtacctgtaacatcgatcatgcatg
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tttgggattggttttgacgcgaaatacaaacattcgcacaaaaccgtcctccatttcaaatgcctacacttgctcactgtatatctct
cttctctcgttttgccacgttgcaGTCTCGTTTCAGCAATCGGAGATACGTACGGTCCCTGCCCTGCTGCT
GCTACACATGCTGACCTCCACCATCAAGCTGACGCTGCTGCCCTACCAGGCAACGAAA
ATCGACGGTGTCAACGTGTACGGATTGACCGTAATCGGATATTTGTGCTACGCGTTGG
CTCAGGTTTCCCTGTTTGGCATCTTTGGCAATCGGCTCATCGAGGAGgtacgtgcgctcggcggtg
ttgccgtgggaagcattctccctgcccatatcgcttcattctcccagatcacacatttgcatacaaaagccagcacacttttgcttcg
ccgtgccatctcggcttctgaatgttttcacttctcccatacttctccgtgcagAGCTCATCCGTGATGAAGCGGC
CTATTCCCTGCCACTGGTACGACGGGTCCGAGGCAAAACCTTCGTCCAGATCGTT
TGTCAGCAGTGCCAGAAAGCGATGACTATTTCCGGAGCCAAAGTTTTCACCGTTTCGC
TCGATCTGTTGCTTCGgtaagtgtagcctggtggctggcacagaacaggctggcacaagaggactttggctctagc
ctgatgggtgtatatgtgtctatttttgtaccattctcgcattctccttccagGTTCTTGAGCCGTTGTCA
CTACTTCATGGTGTGCTGCAGCTGAAGTAAacagccgtggccggaggatgtgttttttcgctcgttcg....
gttgtttgttgccacactttctctggacattttcttactgcaaaaggtttaacaaaacagcaacaaataatccccaaagttttctttt
acagatctttgcaaaaatgattagattttaatagattaacagtgttgaattatctgtcctgtagcaaccgggctgaagaacgttgatt
tggtaaaagtacaaaaggacgttggaattgaaaccacagaaagagtataattatgcacaaagctcaccaagggaatctatgtat
gtgtgatttgcgctcatcaagcacctgtatgtgccttcaactagtcagcaataaagagtacaaatgtttcttagcgccacgtacattg
tcgttcggcgttttaaccgttgttgataatacacaaaagatgataaaaataataataaataatgtttaatatgagtaagtacta...
aatagagaaatcgttttagtatgatcatcacctccaatcattgtttgaaattaactttaactcaaaattaaaccgatgttttact...
ttctgtgagaattattgtggaaagaacttaataatggaagtataatgattgctaactttatgcgtttttcaatttacgaacgtagt...

FIG. 9F

cttcaaacatcgcttcaaaaagtattactaccacattattcatttacttataagttatatatttgccctcttcatcttccatggccagaact
actgcagaaaagcttcttttttgctcgcttccgatggttggtggacgaagtggtaacaaacggcaagcaattagcataaaactatt
ttcgcatcgagatggaaaatgaatgtaccactagaaccgagtgaatgaattacttttcaacttgcacgccaacaccattatctaaag
tacgcacaaacttaaaaaacaaaccccaaatgtcgtccaccttcatccactttcttgctacactttccgaccgagttctgtagcggccag
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gtaccgcaccgcacccgtaccgataccggaaacaaacggtgtgcgcgaagaatccgctagcagccccactggcacgggtatttgctt
ttgggttctgtgttttcttccactgggttgggtgcccctgggctagctcgggctactttcccggggccgcaattttctgcagcccaag
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FIG. 9G

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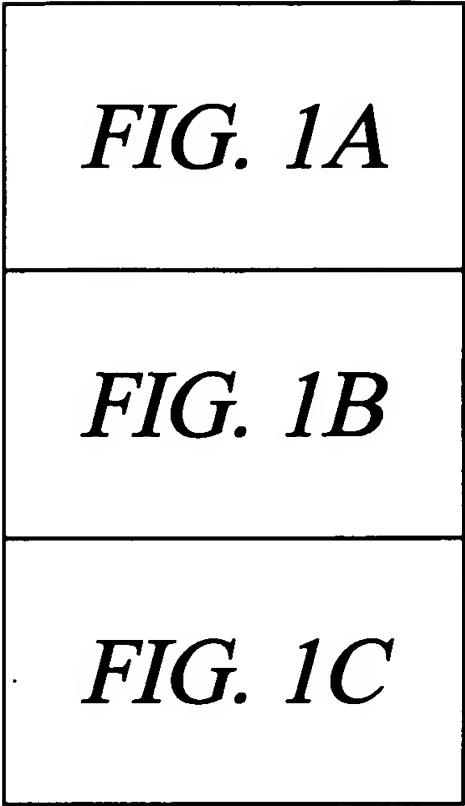


FIG. 1

Now 1A, 1B, 1C
due to
enlargement.

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Anopheles gambiae odorant receptor 1 genomic sequence (SEQ ID NO: 9)

Features:

- 1) Presumed Untranslated 5' and 3' regions are underlined.
- 2) Potential TATA box transcription initiation signal is double underlined.
- 3) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
- 4) Introns are tentatively assigned and are shown in lower case.

Exons are boxed — was highlighted

AGCTTTGTTCATTTATGTTGAAATCTAGCCCATTTTGTATAGTGCTGAACGACGAAACATACGAAAGTACCTCGT
CCGAACACTATCAACATTAATTATACCAAGCTAGAAAGAGATATTATAGTCAAGCCTCAACATCATAGGAAACTTT
AGCAAAACCATTTAATTACATGATGATAAGTCCCACCTCTTACCCAGCACAGGTTTGAGAAGGACGAAAGTATCT
TTACGATAATATTACTCTAAGGTAGTTTGAATAAAATAAAATTACGTGCAAGTGGTGCAATCGGACATCATTC
GAAAGAACTACTAAGTCATACACACACCCAGACGACGTAAGTTTCATCTAGAAAAAACGGTCAGCTCCATC
GAACACGTCAGGACATAACTGCGACATGCGTATGGTCAGTTCACCTAGTGCCAACACACTGGTTCAGGGCACTACCTT
CCGAAGCAGTAGAACCTAATGTATTGGAAATTATTAGGACATACTGCAACATGCATATGGCTAGTTCGCTGGTACC
AACGATGGCACCCAGGACACTATCTGCGGCTTGTAATACTACTGTAAAACTATACAAAAACGGCTTTACCCATACT
TTATCACAAAAACGGCAGGTGAGGGCTGGATTGCTTCAAAGCATTAGAAAAATATAAATTTCAAAGTCCATAATCTCC
TTAAAAGATAGACAaCAGTAGAGAAACACATTTAGTGCTCTTTTCGTTTCGAGTTAGTTGCCTTCTCAAGTAAGCGTTT
AATGCTCAATTGTTGATTCGTTGGATGACTCTCGCTACGTGCTATAGTGGTCAATACTTCCAATTAGATTTCAT
AATTAGTTTCCAATTGTCCACGGAAAACCCaCAAAAGAAAAAACTTGTACTAGGGTGGAATTTTCGAGAACA
ATTGGACACTTCAT**ATG**AAAAAGGACAGCTTTTCAAAATGTTAAATAAACACCGTTGGATCCTTgttggatttca
attctccaaattctgcagaataattctgcaaaattttacaaaactgctcaaccaccaataattccaattaatcatctg

FIG. 1A

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aacattaaaactgataattaagatgagtaattgcttcgtcatcacctaagaaatcgattagtttggataaaaagaa
caaattgaaatacaataaagtcctgaattttatttcgaataaacggcttgaactcatttatttcaaaaaccttttgaga
aattcctcgttgaaaattggtctcctatagttctgctaacgggccacttcaaaagcaagaactaaacaaaatcataat
tatggtgcaagtaactatcagtagtaacagtaatcgccattaaaaactttcctcaatttgcggtcgttacgggctaaa
tacagagcagagtaacgggaagtgatcaacgtcgctattagataacgaggaacgccctccgaagggtgtgtgaagg
acctttcaaatgaaaccaagtactgtttccagttttaaatggatagttataaaatgagccgttcaacgatcggg
catcattgagtttcatcttcgaggagaaatagatcagtgccactgtttaaccgaaagtgaatgaagctgaacaaact
gaaccacggtgggatgcgtacgatcgacgggattcgttctggttcagttgcttgttgaataatttagG CCTAT
GGCCACCGGAAGATACGGATCAGGCAACCGGTACATCGCGTACGGTTGGGCTTTGCGGATCATGTTTCTA
CATCTGTACGCTCTAACGCAAGCCCTATACTTCAAGgATGTGAGGATATTAAATgtgagtcctagttagctattag
tgttccacctgtccataatctgtctttatttgggtagGACATCGCAAATGCATTGTTCTGCTTATGACTCAAGTGA
CGTTGATCTACAAGCTGGAAAAGTTTAACTACAACATCGCACGGATTTCAGGCTTGTCTGCGCAAGCTTAACTGCACA
CTGTATCACCCGAAACAGCGCGAAGAAATTCAGgtaaagcctgctgggaaatatgactaaaaagagtgtacaaacga
ctctcctccaaatgtagCCCCGTTTTACAATCGATGAGTGGAGTGTTTTGCTGATGATCTTTCTCATGTTTGTGGC
TATCTTCACCATCATCATGTGGGTATGTGCCAGCCTTCGACAATGAACGTCTGCCcGTGCCGGCTGGTTCC
CGTGGACTATCACCATTCGGACATAGTGTACGGTGTACTGTTCCTGTATCAAACCATTTGGAAATCGTCATGAGCGCA
ACGTACAACCTTCTCGACCGATACCATGTTTTCCGGCTTGATGCTACACATAAAATGGACAAATTTGTGCGGCTTGGTAG
TATGGTTAAAAAGgtgagttacggcgactacttgccctccagtaaggacaggagttgtttccgttatgatatcatt
ttatcagCTTGGACATGACGTCCCTCCCGAACGCCAATTGGTCGCAACGGATGCGGAATGGAAAGAGATGCGAAAGC
GCATCGACCATCACTCCAAGTGTACGGTACGATGTACGCTAAAGTAACGGAGTGTGTGCTGTTTCACAAGGACATC
TTAAGgtacgaattgggccaattaatgtgtcatttaaaaagcttgacccaactttcacagcttcggcgatgaagt
gcaggacattttccaaagGATCTATCTTCGCGCAAGTATGCGCGTCTGTAAATTATCATTTGTATGACACTGCTGCAAC
TACCGGGGCGATGTTACGATgGCCGATCTGCTGGGCTGTGGGCTCTATTGCTAGTAaAGACATCGCAAGTGTTTA

FIG. 1B

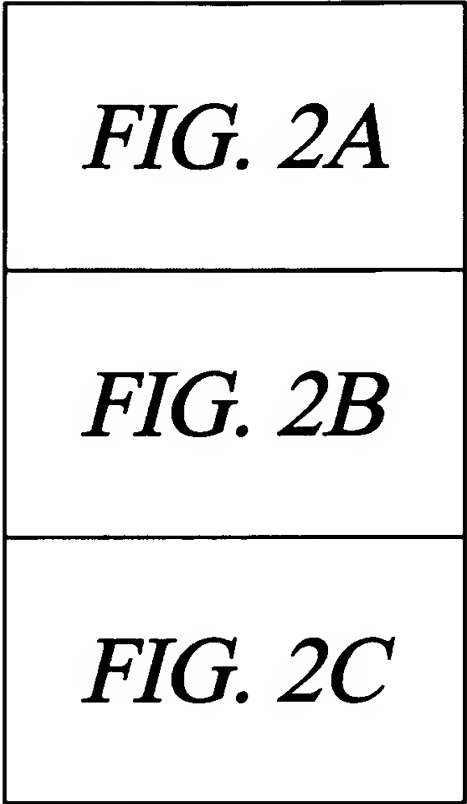
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TTTTCTGTTACGTAGGGAATGAAATCTCCTATACGgtaggttggacacgtagaggaattaaatgtttgggaagaata
tcaataaccaaatagtatgatgttctcgttacagACGGATAAAATTACAGAGTTTGTGGGTTTCCAACTACTTCAAG
TTCGATAAGCGTACCAGCCAAGCAATGATATTTTCTGCAAAAtgtgagatagcgggtgtatttgtcagtcagtaca
ttaaatacgttctctatttcaGACTCTTAAAGATGTTCAACATCAAGGTGGGAAGTGCTTTGAAGGTTACGCTAAAT
CTTCACACATTTTTCAGgtatgtaatatgctgtggtatttagcttgaaataagctacaaactttgaaagtaattt
caatctgtttgtgATTATGAAGCTATCGTACTCCTATCTGGCCGTACTTCAGAGCATGGAATCAGAGTAATGGtG
tTAATATCCtTAA TGTTGAAATTATATTTTGTAGATTTATTGCATAAAAGTAaTaTTTAATTTTATACATCAAACGT
AAGCCCGctaGTTTCAATTAGCCCTTTTCCAAAATTATCAAAATTGATTTCGAATTGATTGCAGAGTTTCAGGAATT
TAATCTGATAGGATATCTTGTATTATCCAATAGAGGTGTGGAAGCGTTCCCAAGCCATTTCGTTTGATAGTTTATAGCA
CCGTCGAGCAGTTGATCGCTGTGATCGCTAGGCGCACCTGATTTTATCTTATCTCGCACCTGTTATGGCAAGGGCG
CTTTTCACACGTTTCACACAATATAATGCACATGTATAATGCATTCTTACTTTAGCATTTTGTACATATAATACC
AAATTATGCATTTTATTCTCACGCAACGATTAGAGGATGACTTcACAAAGGTCCATCTAGTGTAGGAGGTATAC
AATTATACCTCTCAAAATCTCACAGCAtAATGAGAAACAAGGATACCAAGCATACCCTTTTTTTACTTGACAAATT
TCATTTGATTATGTAATAAAGCAGCTGCaCGTCGACTTCCTAAAA

FIG. 1C

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now 2A, 2B, 2C
due to enlargement

FIG. 2

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Anopheles gambiae odorant receptor 2 genomic sequence (SEQ ID NO: 10)

Features:

- 1) Presumed Untranslated 5' and 3' regions are underlined.
- 2) Potential TATA box transcription initiation signal is double underlined.
- 3) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
- 4) Introns are ~~tentatively~~ assigned and are shown in lower case.
- 5) Exons are boxed. — *exs highlighted*

GGGATCCTCTAGAGTCGACCTGCAGGCATGCAAGCTTCCCTCACCGTGACGTGCTAGAAATGGTTCAACATACTCGT
CCGGCAGAGCGAAGACGACGAACAGCGGAATGTCCAGGAAATGTAATGAGATATCACAGCAAGTGAACCCAAACCG
AGCTGTGCGCTTGTGTGCGCTTTAAAAATGGCCCTTCCCTCGCCGCATCTGCTTGGTTTACACGCTTTTCCCAGG
AAATCCACTGACCACTGGCCACACATCAACCACCGGAGCCCTCAGTGCCCGAGCAATATAATTGCTCA
AAAAGTCACGGTACTCAATTAAATTGATTATATAATCAATTTCGTGGCTTCCAACACACCCCTTCTTCCACAATCCATCG
CCGAGTGAGCGAGTATAAAGGTGAAGAAACGTACCTTGCCTTGCCTCACTAAGTGAACCGGATTTCAAAAAGGAACA
TAAACCGCAACCCACAGCCGAAA**ATG**CTGATCGAAGAGTGTCGGATAATTGGTGCAATGTGCGAGTGTGGCTGTTCTC
TGGTCGTATCTGGCGCGCGCGGTGTGCCGCTTCTGGTCGGCTGCATCCCGTGCGCCGTGCTGAACGTTTCCA
GTTCCCTGAAGCTGTACTCGTCCCTGGGCGACATGAGCGAGCTCATCATCAACGGATACTTTACCGTGCTGTACTTTA
ACCTCGTCGt acgtgggcgaggggaggggcaataacctcccacttggtaggataatttcataacctttccatgtgtt
ttttattctctgttgttgccatccagCTCCGAACCTCCTTCTCGTGATCAATCGACGGAAATTGAGACATTTT
TTGAAGGCGTTGCCGCCGAGTACGCTCTCCTCGAGgtaagtcattgggttttctagtttttggggagttgtttaca
ccataaacaccccgacggtaacatttgatcggtcccgcaaaatgtttgtacagAAAAATGACGACATCCGACCCCGT
GCTGGAGCGGTACACACGGCGGGACGCATGCTATCGATAATCGAATCTGTGGCTCGGCCCTTCATTAGTGCCTGCT
TTGTGACCTATCCTCTGTTTGTGCCCGGCGCCTACCGTACGGCGTCAAGATACGGGCGTGACGCTGGCTGGCC
ACCCCGACCTACCAGGTCGTGTTTGTGCTGCAGGTTTACCTTACCTTCCCCCGCTGCTGCATGTACATCCCCTTAC

FIG. 2A

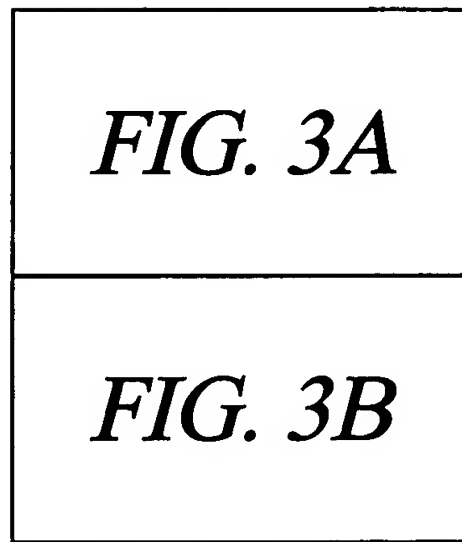
+

FIG. 2B

ttccagtaatccataataaaaaataaagtaaaataagtaaaataatccagtaactgtagtaatac
acaataatctctaagaattaaaattgcatttttgtaatgaaatatgtgattgttcgaatagttcagaaaaacttaaa
aatgcctcagcattaaacagttttgaggttggttcagggcatttagtttagatatatttagtattttaaagcatttggt
ttcattactacaaaaagcaaatttatgagtgaattactttcagttcttctaacgcctatgtgtatgcaattacat
aacaatagctctctttttattgcatttttccttagtaatcctaatactcttcttccctcttgcaATTAAA
GTCCGGCAACGTGTACCCGATGACGTTGGAAATGTTCAAAAATTGCTCAACGTCTCTACTCTCTATTTCACACTGCT
GCGCCGAGTGTAACACTAACTTAACCGGTAAACAAACAAATCCCCTCATCTACTATGCAAGACAGCAAGCAGCC
GATCATCAAAACACCATTAGCAGCCACAAAGTTACCAGCCGCTTATCCACGGGATTGGTGGAAGTTATTGCAC TG
AAGCTCTTTCACCCCAAATTTTCATGGAGGTTCCCTCTCAACCAACCCATTGAAGCGAATAAAAGTATCAGCAACCAG
GCGACGGTGAAAAACGCTGCATTATTGTGCTTGCCTTCAGCATTCAGCGAATGACTCTTAAACTTTTCCATTCAAA
AGTCGCGATGCTACGATACGGAGCGGTGTGTTCGATCCGCCGAGTGCACTCGCAAGCCGGTGATGTGCCGGT
GGAAATGCACAGATCGACACAGCGATAGATAATCGTTTGTTCGCGTAAATGGAGGGGAAAGTAAGCTGCCAGCT
ACTTCATTTCCTATGTTAATTGAAACTCAAGCCAAACGAAACATGCAGAACCCGGTTGGTTGTGTCTCCGCTCCGGGA
AAGGTCTCTGCTCCGGGCATGGATTCTTTCCCTCCGGTGGTTGGGGTATTGTTTAGGTTTATTATTACAAA
TTCATATCCTTCCGCTTCCGCATCAGCCGACCCGGTGCGCCAGACAGATGTGCGGGGCAACAAACTATGC
ACGAACATGGCCAAACACACAGCTTCTATCTGTGTGCTGCACTGTCTCGCTTTCCTGCTGCGTTGCTTGT
GTACTATCATTTAGTCCACGGTTTACTTCTAATTCATTGCACCACGCAAAAGGCTCATCCTTTGCTCGTT
CCGGTTGCAACTTCGACAAAGCGCATGGTTGGGATACGAACAAACCAACTACTCCACCCACTACTACTG
CCACCACCACTAACACACTACACTTGGTTGGGAGCTTGCAGACCCCAAGCAACACGATACAGCTAGCTAGCT
GCTGTGTGCGCTCGAGTCAGCCGACGGTACAAGTTTAACCGGTACAAGCAACTCCCGACCGATCCCAAACTCTG
ACAAGGCACGGCCGATCCGGCAGTACGGTCGGAAACATGGAAATGTTTAATTAAAACTGTAAATTGTCAATCGC
TGCTACAAGTTGTGACACAGGGAGAGAGACAGAGCGCCCGATGGTGATGGTGTAAAGATAGATACAGGAA
AAGAGCGAGAAACATTGGTACGATTGGTGTGGTTAGCAAAATTGATTTCCACTGATTTTGAGTGCAAAATTAAATGC
ATCGAAAAATTGCCATTTCAGGGTAAAGTTGCTCGTGACGGATCCCCGGGCTGCAGGAATTCGATATCAAGCTTAT
CGATACCGTCGACCTCGAGGGGGCCCGGTACCCAGCTTTTGTTCCTTTTAGTGGA

FIG. 2C

+



Now 3A, 3B
due to
enlargement

FIG. 3

+

Anopheles gambiae odorant receptor 3 genomic sequence (SEQ ID NO: 11)

Features:

- 1) Presumed Untranslated 5' and 3' regions are underlined.

2) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.

3) Introns are ~~tentatively~~ assigned and are shown in lower case.

4) Exons are boxed. *was highlighted*

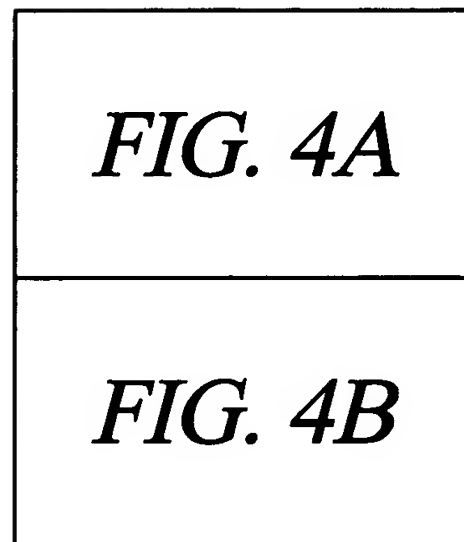
AAGCAGAACACATCAAGAAAGCAATTAGGTGTGTCGTACGTTAGCAAGTAGTTCGCGAGGAGGAATAAAATAG**ATGCC**
TTCTGAGCGGCTTCGCTCTCATTA~~CTT~~CCTTCGGAACTCCTCAAGACAAACGCACGATGGTACTGCCAAAATTAAAGG
ATGAAACAGCAGTGATGCCGTTTCTGCTGCAAAATTCAAACCATTGCCGGACTGTGGGTGACCGTTCCCGAGCGGTAC
CGTTTTATCTCATCTTTTCCCTACTTCTGCGCGATGGTGGTTCTACCCAAAGTGCTGTTCGGTTATCCAGATCTCGA
GGTTCGGGTACGCGGCACGGCCGAGCTGATGTTCGAATCGAACGCATTCTTCGGCATGCTAATGTTTCCTTTCAAC
GCGACAAC**TACGAGCGAT**TGGTGCA**T**CAGCTGCAGGATCTGGCAGCTCTAGgtgagtatgcagccaatcgattgtc
caaaccttcgcaacatccttcgtaacactgctacacttccagTCCCTCCAAGACCTACCCACAGAGCTGGGAGAGTAC
CTGATCTCAGTGAACCGGACGGGTCGATCGGTTCTCCAAATTTACTGCTGCTGTCACTTTTCCATGGCAACGTTCTT
TTGGTT**CATGCCCGTCTGACGACCTAT**TCCGCC**TACTT**GCTGTGGCAACAGCACGCCGGAACCGGT**CGAGCACGTGT**
TGCACCTCGAGGAAGAGCTGTACTTCC**TGAACAT**TCGGACTTCGATGGCGCACTATACGTTTATGTGGCCATTATG
TGGCCACGATCTATACGCTCGGGTTTACC**GGTGGCACA**AGCTGCTGACCA**TTT**CAGCAATGTTAAGTACTGTTC
GGCCATGCTGAAGCTCGTTGCAC**TCCGAATCC**ACTGTCTAGCGAGAGTAGCGCAAGACCGAGCGGAAAGGAGCTGA
ACGAGATTATTTCCATGCATCAGCGGGTACTCAAgtaagtaaatcaaatgaaagtttgcagggaataacttgag
tgtgtctgaccggtgcacatccctagCTGCGGTGTTCC**TGCTGGAGACGACAT**TCCGCTGGGTATTTTTCGTGCAGTTC
ATTCAGTGTA**CAATGATCTGGTGCAGTCTCAT**CCCTCTACATAGCGGTGACCGgtaataagcatttcgctatttcgta
GccttattcaatccatttttgtgaacgtgaatttccccagGGGTT**CAGCTCGACGGTAGCGAATGTATGTGTCCAG**
ATCATTTTGGTGACGGTGAAACTTACGGCTACGGCTACTTCGGAA**CAGATCTAACCAGGAGTGCTTTGG**gtacc

FIG. 3A

ctttggatgaagcttcaaaaagtaattccaaattctgttttcgatttttcccccttttccactagAGCTATGGCGTTG
CCCTCGCCATTTACGATAGCGAGTGGTACAAAGTTTCCATTTCGATGCGCGCAAACTTCGACTGCTACTGCAACGA
TCCCAAAACCGCTCGGCGTAACGGCGGAAAGTTTCGCTTCGTCAATGTGGCCCAAGTTTGGCAAGgtaacattaat
tacagtttgaaaaattctgaagaatgcatcttacttgcccttactgtgttccagATGCTCAAGATGTCCCTATTTCATT
TTACGTAGTACTGAAGGAGCAGTTT**TAG**GAGCTGCTGTTTCCACCCCTGGAAATGGCCCTTTTCGCACTGTCTTCTGT
TTGTTGGACGCACGCAGCACCGAGAGCGCCCTGCACGCACTGACGTAATTTGGCTACTTTTGACGTTTGCACCTTTG
ACAGCTGAAGGACAGGGTACAAATTTTGCTGCTGTATTACGCGCAGCGCATTTGGATACGAAACATTTGGCCACAAG
TTCTACGATTTTAGCGTTTATTACTGTTCGTAGCAGCTTTTCCaCAATAAACACACACAATAACGTACCGACAG
TATTCTTTTCATTGTAGGATAGAGAAAGCCCGGCCAGCAGCCAAACGCAAAACGAAAGCGGCACCCACCG
GGGAAAAACACGGGAGCAAAACGAGAACAGACGAGTAAACAAACCCGGCCGGAACAAACGCGTGCCCGGAA
ACGA

FIG. 3B

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now 4A, 4B
due to
enlargement

FIG. 4

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Anopheles gambiae odorant receptor 4 genomic sequence (SEQ ID NO: 12)

Features:

- 1) Putative Start (ATG) and Stop (TAA) codons are in **BOLD**.
- 2) Introns are tentatively assigned and are shown in lower case.

GGGGAAC TCCCCACCCGACCGACGACGGAAGCTAACGATGTGCAATTGAATAGTCATTAGT
AGCGTTT TTGCTCGCAACGAACTAACCCCTTTGACTTTTAAAGTTCACTACG GTGAGGACAAAA
TCAATAAATTAAATCGAGACCGTTGATGAGCAAAAGAAAAAATAATTTACTGATTTTCATTT
CGTTCCATCGACTACATAATCATAAATTATATGCCACATTTTATATAAGTTT TTGTATCATTTTTA
AACAAACAAAAATGCATCCTTTCGAAATATTAGTCAGGTTGTATCAACA**ATG**AAAGTTTGAACTGT
TTCAAAAATATTCCTCCCGGACACGGTCTTATCCTTCGTCTAAGGCTTTTGCATATCCGTGGGC
ATGAATGGGCAGGATTTTCGGTCTCGCGAATTCGAGTTGGTCATTTTCTGTTCTATTTAATCTT
TCTTGTAATACCGCCACTAACGGGCGGTACACCGATGTCAACGCGTGTACGCACCAAGTGTG
GAATTCCTGTTTAATTGCAATATTACGGCGGCAGTATGTCTTTTGCCCTACGATGTGCCACTTT
CCAAGCGTTCATCCAGGAAC TGAAAGAGCCCTTTCGGTTTGGgtaataattaataaattgcgttattgcat
catcattgttctctttgcagTATGCTCACATTCGTACAGACTAAAGTATAAGCTGACCCGGTTCAACCGTC
GAGCGGATATTATCGCCAAAGTGCAAAACGACCTGCATGGGTGCTGTAAACGCTTTTCTACTGGAT
TGCACCGATACCTTCCATCTGTGCGCACTACTACAGTCCGACCAATTCCACCGAACCCGTGCGG
TTTGTGCAACATTTAGAGGTGAAGTTCTATTGGCTCGAGAAATCGCACCTCAGTCGAGGACTACAT
AACCTTCGTGCTGATCATGCTACCCGTCGTGTTATGTGTGTTACGTATGCAATTTGAAGGTGA
TGACCATCTGCTGCAGCATTTGGACACTGTACACTGTACACCAAGGATGACTATAGAGATGGTAGA
GCAGTTGGAAAGCATGGCATCAGCGGAACGAACTGCCAGCGCCATACGCAACGTGGGCGAGAT
GCACAGTGGTTACTGAAATGCATTAGGCTTTTGAAACACGTCATCCGATCGATGCTGATGCTGC

FIG. 4A

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AGTGGTTGACCTGCCGTGTTAAACTGGAGCATTTCTCTCATCTATCTAACGAACGTGgtagtttgctt
gttggaatccaaacaaagatggctataatggaacttctattacagGCATCTCGCTACAAATCGGTACCGTGGT
GGTAATGTTTTTCTTGCCACTGCCGAAACTTTCCTGTATTGTTACTTGGGACGCGCTTGCGA
CACAAACAGCAGCTGCTGGAGCACGCACTCTATGCTACACGGTGGTACAACTACCCAAATAGCCTT
TCGCAGCAGCATTAGGATGATGTTGAGACAGTCGCAAAAGGCATGCACACATAACGGTGGGAAG
TTTTTTCGCGTTAATTGGAAGAAATTAGCAGGATTGTCAACTTATCCTACTCTGCTTACGTCGT
ACTTAAGGATGTAATAAAGATGGATGTACAG**TGA**ATGTTTMTTMTTGGCTTGGCAACGAATGA
AGTTTTCGGAATCTATATTAGATCTAGAAATTAAATCTAGATGTCATAATATGATCTTGGCCATGA
CCGGTTCCTGGTTTGGAAACCAATTCTCAAAACAATTTTGAACTTAGGGCGAGGCATGAATGTC
CCAAGAACCTATCCAAGTCTGGAACTACATATTACCGAATCTATCCCATTATTGCCCTCGGAAC
GGTTGGTGCTAAATATTGTCCAAATGTTGTCCTGGACCTATCCAGACAAAGATCTTCAATTA
TTCCCTACCACTGGAACTGATTAAATTGATGTAGGAAGTCATGGAGGTGTTACGGGAGAAATTAA
CACTAATGTTCCAACCTCATTTATTTCAAGGCCAATTCTATTATTATATGCCCCTACGGATTGATAC
GTATGTATTACTCCATTTCCTGGACTTTGTCTTATTCTTGCTGCTGATTGGACGTGAATGTTGA
GAAAAGATTCTTATTATGAGTGATACAGAGCCTTTAAATACTCCTACGTTGTTTGCTATTAA
GTATGGCCAGGCTAATCACAAATCGCTACTAATGAACAGAAATCTTCTAATTAAACCTTTCGAT
TGATAGTGTCAATGTCTGAGATAATTGAACCTGCAAAACgATACCTACCTTAACCGAGCAG
AACACATCAAGAAGCAATTAGGTGTGTCGTACGTTAGCAAGTAGTTTCGCGAGGAGGAATAAAAT
AG

FIG. 4B

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+

ANOPHELES GAMBIAE

enlarged

Preferred DNA Codons

| Amino Acids | | | Preferred Codons | | | | | |
|---------------|-----|---|------------------|-----|-----|-----|-----|-----|
| Alanine | Ala | A | GCC | GCG | GCT | GCA | | |
| Cysteine | Cys | C | TGC | TGT | | | | |
| Aspartic acid | Asp | D | GAC | GAT | | | | |
| Glutamic acid | Glu | E | GAG | GAA | | | | |
| Phenylalanine | Phe | F | TTC | TTT | | | | |
| Glycine | Gly | G | GGC | GGT | GGA | GGG | | |
| Histidine | His | H | CAC | CAT | | | | |
| Isoleucine | Ile | I | ATC | ATT | ATA | | | |
| Lysine | Lys | K | AAG | AAA | | | | |
| Leucine | Leu | L | CTG | CTC | TTG | CTT | CTA | TTA |
| Methionine | Met | M | ATG | | | | | |
| Asparagine | Asn | N | AAC | AAT | | | | |
| Proline | Pro | P | CCG | CCC | CCA | CCT | | |
| Glutamine | Gln | Q | CAG | CAA | | | | |
| Arginine | Arg | R | CGC | CGG | CGT | CGA | AGA | AGG |
| Serine | Ser | S | TCG | AGC | TCC | AGT | TCT | TCA |
| Threonine | Thr | T | ACG | ACC | ACT | ACA | | |
| Valine | Val | V | GTG | GTC | GTT | GTA | | |
| Tryptophan | Trp | W | TGG | | | | | |
| Tyrosine | Tyr | Y | TAC | TAT | | | | |

[http://www.kazusa.or.jp/codon/cgi-bin/showcodon\(con'd on next line\)
.cgi?species=Anopheles+gambiae+\[gbinv\]](http://www.kazusa.or.jp/codon/cgi-bin/showcodon(con'd on next line).cgi?species=Anopheles+gambiae+[gbinv])

FIG. 5

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+

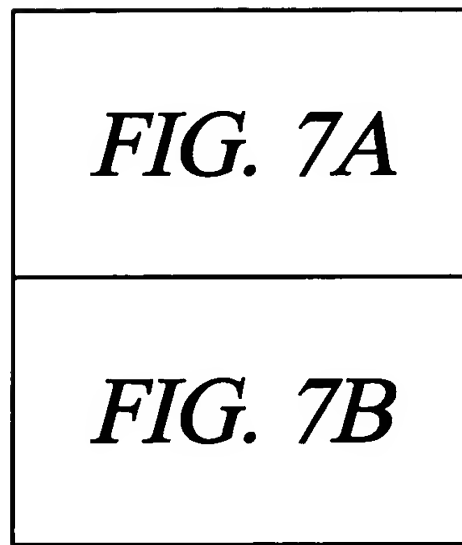
enlarged

| Name | SEQ ID NO |
|----------------------------------|---------------|
| Arrestin 1 (cDNA) | SEQ ID NO: 1 |
| Arrestin 1 (polypeptide) | SEQ ID NO: 2 |
| Odorant Receptor 1 (cDNA) | SEQ ID NO: 3 |
| Odorant Receptor 1 (polypeptide) | SEQ ID NO: 4 |
| Odorant Receptor 2 (cDNA) | SEQ ID NO: 5 |
| Odorant Receptor 2 (polypeptide) | SEQ ID NO: 6 |
| Odorant Receptor 3 (cDNA) | SEQ ID NO: 7 |
| Odorant Receptor 3 (polypeptide) | SEQ ID NO: 8 |
| Odorant Receptor 4 (cDNA) | SEQ ID NO: 13 |
| Odorant Receptor 4 (polypeptide) | SEQ ID NO: 14 |
| Odorant Receptor 5 (cDNA) | SEQ ID NO: 15 |
| Odorant Receptor 5 (polypeptide) | SEQ ID NO: 16 |
| Odorant Receptor 6 (cDNA) | SEQ ID NO: 17 |
| Odorant Receptor 6 (polypeptide) | SEQ ID NO: 18 |
| Odorant Receptor 7 (cDNA) | SEQ ID NO: 19 |
| Odorant Receptor 7 (polypeptide) | SEQ ID NO: 20 |

FIG. 6

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Now 7A, 7B
due to
enlargement

FIG. 7

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Anopheles gambiae odorant receptor 5 genomic sequence (SEQ ID NO: 21)

Predicted Exons: ITALICIZED, UNDERLINED AND BOXED

Introns: lowercase.

was 3/19/10

tctagacttgaacccatgacgggcattttattgagtcgttcgagtgactgtaccacgggaccaccgctttatcactatcactatt
aattaattataatatgctttttagcagcctaccgggttttctctggatatcttaagtcccatattgattatcaagatagaa
caacaacttgaccttaataatacattacgtacccttaatacaacctgtgcatcaaggagtttcggaagcaaaaatccgattgtct
gatgtgttgattccatccgattcgttactggttctgcaaaatcgccaataatacggcaatgtccttattcgaatgattgaatcaacat
cacattgttgcatctcgtttttgctgcaaatatgttattgcaaaaggcaagtaatgtgcttaagagtaataacaattcgcctg
tccattttttgccaccagtggtccagaacccgtgccttttagtccttcgaatacatccgaccagtcagcaagtgcatd[ATGG]
TGCTACCGAAGCTGTCCGAACCGTACGCCGTGATGCCGCTTCTACTACGCCCTGCAGCG
TTTCGGTGGGCTGTGGGTGAACGACGCTATCGCTACAAGTTCCGGTTGGCATTTTA
AGCTTCTGTCTAGTAGTATTCGGAAGGTGGCTTCCGGCTATCCAGATTTAGAGAC
AATGGTTCGGGAACAGCTGAGCTGATTTTCGAATGGAACGTA CTGTTTGGGATGTTG
CTGTTTTCTCTCAAGCTAGACGACTATGATGATCTGGGTACCCGGTACAAGGACATATC
AAAGATTGgtgctgataatgattgataaaaggaaacctttgagcaactcctatcccttcaag[CTTTCCGTAAGGAC
GTTCCCTCGCAGATGGGCGACTATCTGGTACGGCATCAATCATCGTATCGGTTTTC
CAAGATCTACTGCTGCAGCCCATCTGTGTTTGGCCATCTTCTACTGGGTGGCTCCTTCGT
CCAGCACCTACCTAGCCTAGCCTGGGGCACGAAACAGATCCGTCCTCCGGAACATGT
GCTACACCTGGAGGAGCTGTACTGGTTTCACACCCGGCTCTCGCTGTAGATTAC
TCCATATTCAACGCCCATCATGCTGCCCTACAATCTTTATGCTAGCGTACTTCGGTGGACT
AAAGCTGCTAACCATCTTCAGCAACGTGAAGTACTGTTCGGCAATGCTCAGGCTTGTG

FIG. 7A

+

GCGATGAGAAATCCAGTTCATGGACCGGCTGGACGAGCGGAAAGGAACTGA
TCGAAATCATCGTCATGCATCAGAGGCGCTAAAGgtaaagggtgtccgggtatgttgtagaatacat
ctagctgctttcagATGTGTGGAGCTGTTGGAAATCATCTTTCGGTGGGTTTCTGGGACAG
TTCATACAGTGCGTAATGATCTGCTGTCAGCTTGGTTCTGTACGTCGCCGTTACCGgtaacta
aaagcactgtagtgatctgtctgccacaccattcactgctgtgtctgtttgtcactcttccagGGTCTCAGCACAAAG
CGGCAAAACGTGGGTGTACTGCTAACAGTGGAAACCTACGGATTCTGCTA
CTTGGCAGTGATCTTACCCTCGGAGGCAAGTTGTTATTGCTGAgtttcagttacttttccgttcccc
tctaaccgtaccactgtaccattgtttgagacagagcttgagcgtagCACGTGCTGCGTACGGTAGCCTCTGG
TATCGCCGTTCCGGTTCCGATTCAACGGAAAGCTTCCGAATGGTACTGCAGCGTGCCCAGA
AACCGGTCGGCATCTCGGCTGGGAAGTTTTGCTTCGACATTGAGCAGTTTGGCAA
TgtatggggagaccttccactgtggcaagaagattttctttattaatgcattctttaattacagATGGCAAAACATCA
TACTCGTTCTACATCGTTCTGAAGGATCAATTTTAAaggggaactccccaccaccagacgagaa
agctaacgatgtgcaattgaatagtcattagtagcgtttttgctcgaaacgaaactaaccctttgactttttaagttaactacggtgag
gacaaaaatcaataaataatcgagaccggttgatgagcaaaagaaaaataattttactgattttcatttcgttccatcgacta
cataatcataattatatgccacattttattataagtttttg

FIG. 7B

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enlarged

Anopheles gambiae odorant receptor 6 partial genomic sequence (SEQ ID NO: 22)

These are the predicted last three exons of another candidate *Anopheles gambiae* odorant receptor.

was high/ish

Predicted Exons: ITALICIZED, UNDERLINED AND BOXED
Introns: lowercase.

aacacccattatcgcaaaattagattaccggttgaaagggttcccttcctggctgtttctcactctctctctctctctta
ttgatgccgtatgccggtctatagctagTTATGCTTACCGGATGTTGCCGATCGCGCACGGTGCTTT
TCCGCATACGCCAGTGACACTTGATGGCGGTGGTGATGACGCTCTGCTGCGCACCGTT
TTCTGCTCGTGAGTCAGACCCTTTTCATTTCCTGCAATATCCTGTTTCTTTCCCGACCC
ACAGACGGTTAGACGGATATATGCTGGTAAAGTTTGTCTCTTCATGCTGTGCTTTCTG
ATCGAGCTGCTGATGCTGTGTGCGGTACGGTGAGGATATTGTGGAAATCCgtaaggcaccaggc
ggtgatgagcgcgagtaattgaagcttttgcctttaaacaacatcagagCCTTGGGGTGATTGATGCCGCT
TACGGTTGCCGAATGGTACCGGGAAGGTCGGTGCGGTTCCATCGATCCGCTGCTGCAAA
TTATACACCGCAGCCAGTCCTCATACTGACCGCATGGAAATTTGGCCCATCCAA
ATGAGTACTTTCAGTCAGgtgagttgccaatgattgccgttgcgttaataattcagtaagagtgctcttcccttag
ATCCTGCAAGCTTCCTGGTCCCTACTTACCCCTCCTGAAGACCGTCTACGGGAATAAgtaa
gctgagagagagagagcagtagcttcccttggatgaatcaatagatttctaatacgaaccattgaaaaatgaatca
acatttcgtagtgcacaatattgtaccattctatacagcttcaccagcgaagcgtttgttcacaggaaccaaacggttcga
caagccgcgtcacctgctggc

FIG. 8

+

+

FIG. 9A

FIG. 9B

FIG. 9C

FIG. 9D

FIG. 9E

FIG. 9F

FIG. 9G

Now 9A, 9B, 9C
9D, 9E, ~~9F~~, 9G
due to
enlargement

FIG. 9

+

was highlighted

BOXED

- was highlighted

ccgccggcagggtgacttacgggtctgacttgctggctgcttgttacggcaaacggctacacaaaggcaatcgaattattttcc
tatcacgctgccttacaggccctgctggtaggcaaaagaatgtgcaaaagtcttcatttggttgctgcttgctgtaacgtgtg
gcacggttgcatcgctaaggtttcgggtgtgagccgagaaagtgtgcagatcgaaatctcttgtgtgtgtgtgtgcagtgggaa
gcattgtttagtgaagtgaaaagaatgtgtaaaatgcaagtcacggaccgaagtagctcgccctcgttgccgacct
gatgccgaacattcgggttgatgcaggccaggctcaactttctgttcggctacgtcacggcccgatatactgacccgaaggtgtac
tcctggtgacgctgcctATGGTGTCTGATCCAGTTCTTCGCCATCCTCGGCAACCTGGCGACCGA
ACGCCGACGACGTGAACGAGCTGACCGCCAACACGATCACGACCCCTGTCTTCACGCA
CTCGGTCACCAAGTTCATCTACTTTCGGGTCAACTCGGAGAACTTCTACCGGACGCTC
GCCATCTGGAACCAGACCAACACGACCCGCTGTTTGCCGAATCGGACGCCCGGTACC
ATTCGATTCGCTCGCCAAGATGCGGAAGCTGCTGCTGTGATGGCCACCACCGT
CCTGTCGGTGTCCgtatgtgtgtatgtgtggccgttgggaaagtgtcttggcgagaacccaatctactgttacgc
ttgactgggtttttgtttttctcggtagggacgggataaaatatctgaaagaataattgagtcacccacagggatgcaag
acatgcaggcagagatttgggttgattatcacccgacacgaatatcttcaggttcataagcttcaccggtgaaaaagga
actccccattccctgtttttttttctctctcgataaaattactcatcgcttttctgtttttttttgtgtgtgttcttcttcttcac

cctactagCCTGGGTACGATAACATTTTCGGCGAGAGCGTCAAGACTGTGCTCGATAAG
GCAACCAACGAGACGTACACGCTGATATACCCGGCTGCCCATCAAGTCCTGCTATC
CGTGGAATGCAATGAGCGGACCGGTACATTTCTCTTCTATCTACCAGTACGTTG
GCGGAATgtcctgcggtcacagttggcagtcagtgagcggaacacggcgaaataatgggactaaacgggtcttcacaga
gccaacacattcctacagcaattgcataccttcggcggtcggactgggcaatgcagtacaacatcctcgccataaagttagcaat
tcgagcgacaaatgttgccgtgttagggctttttgtgataatagtcgtttttgtcctcctcgttatcaaaactctatcaacggaggaaa
tcattttcgtcaaatgcctacagctcaagttcaaggtaatcgaggggtgggatcaactttttattcattttgctaacgccccca
tcaacaaattctatgttctcaatggcaagattactgcccccaacgaacggcaaaagaaagcgacgattatga
agatgtccaaaccattgccccgcgaccttatctgatgttggggatggctttactgtctgtactttcaggcacaaaaggaa
atgaaccagcgagctcgtttgccggttcggagggttcttcaggcactgagctgagtaataatcgaaacgatttttacgattc
tggtaccagtttatgatgtggcctgcattacagtgggcaattataacctgatgttcatttcattgttaagttgtgctggaacg
ccgtaacgattaattcttttcaaaagagattctttcaaaagagattcaaaatgtgtatacaaaatgtaacgaatggaccgtactgg
aggttgcggaagtaacgttttaaaatattcatcacaatcctctgcaaaactgtgcttaattaattggtgcacaataagtttaaac
gtggcggcagatgtgtcgttcgcttccttcctccagcaagctcgtgcgaaataatttattccatcattttaataacagccgtttgtg
cattttaattagcaaaagcaataataaaagcagctaaccatccccattaaacaaagtcttcgggccccaatgttatggcggtgga
aagtaatggttttaccagtggaagtgcctttcccatcgtgggtacttcgcgatatcttcttatacaagtgcatacagaaaaaa
ggacaaatcctccttgctatgggtctaaggccagcttcggtagccggttcggtatgataaagtttgatgggtttttaacatt
actccgctctaaccacctaattggacttttcagtgaactaaagtaaacgaccagcggtacgaccgagccacggttgatt

FIG. 9B

FIG. 9C

FIG. 9D

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+

ttttgcagcgggtatagtaaattgcatactttaaggcgtgatttcaaatgtagcgttccgtatgcagaaacgccaatggattatgc
aatttaacaatgctgcttccttaacattcaataaacggcttattaaggaaactttttgtgcaattttgtttaacagcaaatagtttagc
tcagaacgatcacatttagtatcgcttcaacaagaactcttttaaacacacacaatttgtaatgccattccctcgagaaagtttcttgctc
agtcctcctctgcatacacagcaacaacaacctgctcatgtttcctgctcgtttcctagctgttttgaaacgttattccgattcctgtgct
tgcccgcctttcttacaatacacacaatgggttcagatttcgtcttatttgaccactgctttcgtgctgaagcccgtggaaacaa
tgcgccaagctcagcatccagcccatgtataaatgagccacgcgacagatttttagacatcgctttcgctctgcacccggagggtggtt
ttattcttggttccgattcccacgtccattcgctgggtccgtccggccggcgaacccgtaagccgtgcggggaattacgcaatcga
aacgagccagaaaatgagcacgcgcaaaatgcaaaagaaaatcccccttttgagtgtgctcctgccaccactcatctccccaaactggtgg
gtgaaaaaacctgtgccccctctcttccagaaaaaacgcctcgctcgacaaaaaacatgctcgggtgaagctgcgtatgt
cgcagaagctcaaaacacgcgccagcaagcatcaacaatttctatcaaacacccacagcgcgaacccgggtgcactgta
ctcagtagcgaagatgctcagattgtcccgtgcgtcttcgatgccgtttcggagcgggaagccatcgcttgccaacgttggcgat
gtcttttagccgtgatttgaaattttctgaatatcacaggcggcggttgccctgcaagggttgcttcccacacgagcattgcttt
ccgtaccgcggtgggcgagttttcaacgcaaccttctacaagcaacgcccaacgcctgggagcgatatttaacagaaacaagaa
catcccgaaacttcagcacatgccgtgatttgccctgttggaataagcttttgtagcgtgtgagttgaacgagctctattttccagcgat
gggtggcatttgtgtggcatgctatcgctcagcttttcttgaaatctttacctcctccatcgctccattagtagcacgcgtatggaaaaatgg
gtgcaacggatcagaacggattttccgcgacagacttaataaagggaagcaacgcgttttttgcatgtgtgttttatgagcttt
atgccgttactttgcaattaaaaatagcaaaaaataaacagtttttttgtaagcggattacaagaatgtatcagaatatattacgtg
aaacattcattcatgtctgttaacgctcaaatagatgttttgtaacacggattgcataaccttgccggtatcggttacattttcgccta
acagtatgcaatctgttagctttgttgaatgactgcgttggttagtacaatatatttacaccgcgtaatttatctcacaatatgc
aaaaaatgtcaatctgtatcgattattcacacaaatcagatcccggaaccagtgtagcccaatgtgctcttattgaattaccacga
acaaatcaacctgatccccgggtccgttggaacaacagcttgccggcgaagccgctcaggtttcgtgcactaccgtgctgccattttgct
gccctcatcgaacagataaacagaaggcaactctgtgagcatcgcaatgcccgctcgaaagtccgtcgaaaaatgggcctaattc
aatttgacgcatttaccgcgaacaattgcgcgaaggctgtcaagtggttccacgaactgcgacaacaagcacacacaacaacac

FIG. 9E

aatgttatcgttttcggcatgtttctcggtaacaagcgtgtggcgctatgtggcatgccgattcccagacagagtgatcgatagtaaa
ttagcctatccggtagcattcaatttcctttctatcctcgcaacaagcccattctggggaggcgtggtgaagctttcaaggcat
tgtgaacaaatgtcctggttcggagggtatgtgggaaagcaaacacggtgcccatcgtctaccgtcaatcgatcatgcatg
atgtattaatattgtgttattcacctcggtatctatcgctcgctgtgttcggatttcgggaagtcaaggaaaaagcgactcca
tttgggattggttttgcagcgaaatacaaaacattcgcacaaaacgctcctcatttcaaatgcctacacttgctactgtatatctct
ctttctctcgttttggcacgttgcaagTCTCGTTTCAGCAATCGGAGATACGTACGGTCTCCTGCCCTGCTGCT
GCTACACATGCTGACCTCCACCATCAAGCTGACGCTGCTCGCTTACCAGCAACGAAA
ATCGACGGTGTCAACGTGTACGGATTGACCGTAATCGGATATTGTGCTACGCGTTGG
CTCAGGTTTTCCTGTTTGGCATCTTTGGCAATCGGCTCATCGAGGAGgtacgtgcgctcggcgtg
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CTATTCCCTGCCACTGGTACGACGGGTCCGAGGCAAAACCTTCGTCCAGATCGTT
TGTCAGCAGTGCCAGAGGCGATGACTATTTCGGAGCCAAAGTTTTCACCGTTTCGC
TCGATCTGTTGCTTCGgtaaagttagcctggtggcgacagaaacaggctggcaaaacaggacttggctctagc
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CTACTTCATGCTGCTGCTGCTGAAGTAAacagccgtggccggaggatgtgttttttcgctcgttcg....
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tggtaaaagtaacaaaagggaagttggaaaattggaaccacagaaagagtataatttatgcaaaagctcaccaagggaatactatgtat
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tcgtttcggcgttttaaccgttgttgataatacacaaaagatgataaaaataaataaataatgtaataatgagtaagtaacta
aatagagaaaatcgttttagtatgatcatcctccaatcatttgtttgaaaattaactttaactcaaaataaaccgatgttttact
ttctgtgagaattattgtggaagaacttaataatggaagtataatgattgtaacttttatgcgtttttcaatttacgaacgtagt...

FIG. 9F

FIG. 9G